

Serial Number: 09/687,860CRF Processing Date: 10/27/2000Edited by: 10Verified by: 10 (STIC staff)**ENTERED**

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: \_\_\_\_\_
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other \_\_\_\_\_
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: \_\_\_\_\_
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: \_\_\_\_\_
- ☒ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: 22
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: \_\_\_\_\_
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: \_\_\_\_\_
- ☐ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as \_\_\_\_\_
- ☐ Inserted mandatory headings, specifically: \_\_\_\_\_
- ☐ Corrected an obvious error in the response, specifically: \_\_\_\_\_
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: \_\_\_\_\_
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted **ending** stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: \_\_\_\_\_
- ☐ Other: \_\_\_\_\_

\*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

OIPE

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/687,860

DATE: 10/27/2000  
TIME: 13:08:59

Input Set : A:\36737 sequence listing.txt  
Output Set: N:\CRF3\10272000\I687860.raw

Does Not Comply  
Corrected Diskette Needed

3 <110> APPLICANT: Asundi, Vinod  
4 Ford, John E.  
5 Drmanac, Radoje T.  
6 Liu, Chenghua  
7 Tang, Y. Tom  
8 Yamasaki, Vicky  
9 Yeung, George  
10 Zhang, Jie  
11 Zhou, Ping  
13 <120> TITLE OF INVENTION: EGF MOTIF PROTEIN, EGFL6, MATERIALS AND METHODS  
15 <130> FILE REFERENCE: 28110/36737  
C--> 17 <140> CURRENT APPLICATION NUMBER: US/09/687,860  
C--> 17 <141> CURRENT FILING DATE: 2000-10-13  
17 <150> PRIOR APPLICATION NUMBER: US 09/620,312  
18 <151> PRIOR FILING DATE: 2000-07-19  
20 <150> PRIOR APPLICATION NUMBER: US 09/363,316  
21 <151> PRIOR FILING DATE: 1999-07-28  
23 <160> NUMBER OF SEQ ID NOS: 32  
25 <170> SOFTWARE: FastSEQ for Windows Version 3.0

## ERRORLED SEQUENCES

841 <210> SEQ ID NO: 22  
842 <211> LENGTH: 20  
843 <212> TYPE: DNA  
844 <213> ORGANISM: Artificial Sequence  
846 <220> FEATURE:  
847 <223> OTHER INFORMATION: Description of Artificial Sequence: primer  
849 <400> SEQUENCE: 22  
E--> 850 gggaaactgac ata~~caaa~~agtc

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## VERIFICATION SUMMARY

DATE: 10/27/2000

PATENT APPLICATION: US/09/687,860

TIME: 13:09:01

Input Set : A:\36737 sequence listing.txt

Output Set: N:\CRF3\10272000\I687860.raw

L:17 M:270 C: Current Application Number differs, Replaced Current Application No  
L:17 M:271 C: Current Filing Date differs, Replaced Current Filing Date  
L:61 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1  
L:62 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1  
L:103 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2  
L:104 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2  
L:206 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2  
L:207 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2  
L:308 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4  
L:416 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:5  
L:416 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:5  
L:535 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6  
L:574 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7  
L:576 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7  
L:578 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7  
L:592 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8  
L:594 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8  
L:596 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8  
L:639 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10  
L:641 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10  
L:643 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10  
L:661 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11  
L:805 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18  
L:850 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:20 SEQ:22  
L:1186 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27  
L:1198 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27  
L:1203 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27  
L:1207 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27  
L:1211 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27  
L:1215 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27  
L:1219 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27  
L:1223 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27  
L:1227 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27  
L:1231 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27  
L:1235 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27  
L:1239 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27  
L:1243 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27  
L:1247 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27  
L:1251 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27  
L:1255 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27  
L:1259 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27  
L:1263 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27  
L:1267 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27  
L:1271 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27  
L:1275 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27  
L:1279 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27  
L:1283 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27  
L:1287 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27

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L:1291 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27  
L:1295 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27  
L:1299 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27  
L:1303 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27  
L:1307 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27  
L:1311 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27  
L:1315 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27  
L:1319 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27  
L:1323 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27  
L:1327 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27  
L:1331 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27  
L:1335 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:27  
L:1482 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29  
L:1494 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:29  
L:1498 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:29  
L:1502 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:29  
L:1506 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:29  
L:1510 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:29  
L:1514 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:29  
L:1518 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:29  
L:1522 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:29  
L:1526 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:29  
L:1530 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:29  
L:1534 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:29  
L:1538 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:29  
L:1542 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:29  
L:1546 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:29  
L:1550 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:29